Assessing the genetic diversity of Cu resistance in mine tailings through high-throughput

recovery of full length copA genes

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Running title: Metagenomics for copA diversity in mine tailings

16		84	455
GMTCASCSA	R		VLVVACPCA-
GMTCASCSSI	~	CASCVGRV	VLVVACPCA-
GMTCASCSSI		CASCVGRV	VLVVACPCA-
GMSCASCVG	~~	CASCVGRV	VLIIACPCA-
GMGCAACAG	R		VLIIACPCA-
GMTCAMCVK	s		VLVVACPCA-
GMTCASCAG	R		VLVIACPCS-
GMTCANCSA	R		VLVIACPCA-
GMTCAACSN	R GMT	CAACSNRI	VLVIACPCS-
GMTCAACAN		CAACANRV	VLVIACPCA-
GMTCAACAA	R GMT	CAACANRI	VLVIACPCA-
GHACHHE			VLIIACPCA-
NMNCAGCVA			VLIIACPCA-
GLSCGHCVK		CASCVSRV	VLIIACPCA-
GLSCGHCVK		CASCVTRV	VLIIACPCA-
GLSCGHCVA	The state of the s	CASCVSKV	VLIIACPCA-
GLSCGHCVA	The state of	CASCVSKV	VLIIACPCA- VLVVACPCA-
GMKCAGCVA: HARHQEHGM			VMVIACPHA-
HOEHGM			VMVIACPHA-
HSGHQK			VMVIACPHA-
CYHCGL	The second second	CAACGWLI	LLVATCPCA-
CYHCGL		CAACGWLI	MLVATCPCA-
VN			LLVIGCPGA-
WVRCDSRRAY	v		MLLIACPCA-
GMACAFCAS!	r		VSTIAYPCA-
			FMVVASPCA-
LHWHGL	PHQH	AHCHL	
	YHŜH	FHCHD	
	MHSH	NHCHN	
	YHAH	FHCHH	
	YHSH	FHCHQ	
	YHSH	FHCHL	
	YHSH	YHCHL	
	YHSH	YHCHM	
	YHSH	YHCHM	
100	142	570	
		-	